

## SEQUENCE LISTING

<110> Havenga, Menzo

<120> Chimeric Adenoviruses

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<140>09/348,354

<141> 1999-07-07

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<170> PatentIn Ver. 2.1

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Gly Tyr Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Pro Pro Phe Val 35 40 45

Ser Ser Asn Gly Phe Gln Asn Phe Pro Pro Gly Val Leu Ser Leu Lys 50 55 60

Leu Ala Asp Pro Ile Thr Ile Asn Asn Gln Asn Val Ser Leu Lys Val 65 70 75 80

Gly Gly Leu Thr Leu Gln Glu Glu Thr Gly Lys Leu Thr Val Asn 85 90 95

Thr Glu Pro Pro Leu His Leu Thr Asn Asn Lys Leu Gly Ile Ala Leu 100 105 110

Asp Ala Pro Phe Asp Val Ile Asp Asn Lys Leu Thr Leu Leu Ala Gly 115 120 125

His Gly Leu Ser Ile Ile Thr Lys Glu Thr Ser Thr Leu Pro Gly Leu 130 135 140

Val Asn Thr Leu Val Val Leu Thr Gly Lys Gly Ile Gly Thr Asp Leu 145 150 155 160

Ser Asn Asn Gly Gly Asn Ile Cys Val Arg Val Gly Glu Gly Gly 165 170 175

Leu Ser Phe Asn Asp Asn Gly Asp Leu Val Ala Phe Asn Lys Lys Glu
180 185 190

Asp Lys Arg Thr Leu Trp Thr Thr Pro Asp Thr Ser Pro Asn Cys Arg
195 200 205

Ile Asp Gln Asp Lys Asp Ser Lys Leu Ser Leu Val Leu Thr Lys Cys 210 215 220

Gly Ser Gln Ile Leu Ala Asn Val Ser Leu Ile Val Val Ala Gly Arg 225 230 235 240
Tyr Lys Ile Ile Asn Asn Asn Thr Asn Pro Ala Leu Lys Gly Phe Thr 245 250 255
Ile Lys Leu Leu Phe Asp Lys Asn Gly Val Leu Met Glu Ser Ser Asn 260 265 270
Leu Gly Lys Ser Tyr Trp Asn Phe Arg Asn Gln Asn Ser Ile Met Ser 275 280 285
Thr Ala Tyr Glu Lys Ala Ile Gly Phe Met Pro Asn Leu Val Ala Tyr 290 295 300
Pro Lys Pro Thr Thr Gly Ser Lys Lys Tyr Ala Arg Asp Ile Val Tyr 305 310 315 320
Gly Asn Ile Tyr Leu Gly Gly Lys Pro His Gln Pro Val Thr Ile Lys 325 330 335
Thr Thr Phe Asn Gln Glu Thr Gly Cys Glu Tyr Ser Ile Thr Phe Asp 340 345 350
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Gly Tyr Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Pro Pro Phe Val 35 40 45

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Gly Lys Ser Tyr Trp Asn Phe Arg Asn Glu Asn Ser Ile Met Ser Thr

Ala Tyr Glu Lys Ala Ile Gly Phe Met Pro Asn Leu Val Ala Tyr Pro Lys Pro Thr Ala Gly Ser Lys Lys Tyr Ala Arg Asp Ile Val Tyr Gly Asn Ile Tyr Leu Gly Gly Lys Pro Asp Gln Pro Val Thr Ile Lys Thr Thr Phe Asn Glu Thr Gly Cys Glu Tyr Ser Ile Thr Phe Asp Phe Ser Trp Ala Lys Thr Tyr Val Asn Val Glu Phe Glu Thr Thr Ser Phe Thr Phe Ser Tyr Ile Ala Gln Glu <210> 16 <211>391 <212> PRT <213> Human Adenovirus 13 Fiber Protein <400> 16 Xaa Xaa Xaa Xaa Ser Ala Pro Thr Ile Phe Met Leu Leu Gln Met Lys Arg Ala Arg Ser Ser Xaa Asp Thr Phe Asn Pro Val Tyr Pro Tyr Gly Tyr Ala Arg Asn Gln Asn Ile Xaa Phe Xaa Thr Pro Pro Phe Val Xaa Ser Asp Gly Phe Lys Asn Phe Pro Pro Gly Val Leu Ser Leu Lys Leu Ala Asp Pro Ile Thr Ile Ala Asn Gly Asp Val Ser Leu Lys Val Gly Gly Leu Thr Leu Gln Glu Gly Ser Leu Thr Val Asp Pro Lys Ala Pro Leu Gln Leu Ala Asn Asp Lys Lys Leu Glu Leu Val Tyr Asp 

Asp Pro Phe Glu V 115		Lys Leu Ser 125	Leu Lys Val Gly His
Gly Leu Lys Val L 130 13	_		Gly Leu Lys Asp Leu
Ile Gly Lys Leu Va 145 150	l Val Leu Thr ( 155	Gly Lys Gly I 160	le Gly Ile Glu Asn
Leu Gln Asn Asp A	Asp Gly Ser Ser 170	r Arg Gly Val 175	Gly Ile Asn Val Arg
Leu Gly Thr Asp C 180	Gly Leu Ser 185	Phe Asp Arg	Lys Gly Glu Leu Val
Ala Trp Asn Arg L 195		g Arg Thr Le 205	u Trp Thr Thr Pro Asp
Pro Ser Pro Asn C			Asp Ser Lys Leu Thr
Leu Val Leu Thr L 225 230	ys Cys Gly Ser 235	Gln Ile Leu A 240	Ala Thr Val Ser Ile
Ile Val Leu Lys Gl	y Lys Tyr Glu l 250	Phe Val Lys I 255	ys Glu Thr Glu Pro
Lys Ser Phe Asp V 260	al Lys Leu Leu 265	Phe Asp Ser 270	Lys Gly Val Leu Leu
Pro Thr Ser Asn Lo 275		Tyr Trp Asn 285	Tyr Arg Ser Tyr Asp
Asn Asn Ile Gly Tl			Pro Phe Met Pro Asn
Leu Lys Ala Tyr P 305 310	ro Lys Pro Thr 315	Lys Thr Ala S 320	Ser Asp Lys Ala Glu
Asn Lys Ile Ser Se 325	r Ala Lys Asn l 330	Lys Ile Val Se 335	er Asn Phe Tyr Phe
Gly Gly Gln Ala T 340	yr Gln Pro Gly 345	Thr Ile Ile Ile	Lys Phe Asn Glu

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Leu Thr Thr Tyr Arg Asn Ile Asn Phe Thr Ala Glu Leu Phe Phe Asp 165 170 175
Ser Ala Gly Asn Leu Leu Thr Ser Leu Ser Ser Leu Lys Thr Pro Leu 180 185 190
Asn His Lys Ser Gly Gln Thr Trp Leu Leu Val Pro Leu Leu Met Leu 195 200 205
Lys Val Ser Cys Pro Ala Gln Leu Leu Ile Leu Ser Ile Ile Ile Leu 210 215 220
Glu Lys Asn Lys Thr Thr Phe Thr Glu Leu Val Thr Thr Gln Leu Val 225 230 235 240
Ile Thr Leu Leu Phe Pro Leu Thr Ile Ser Val Met Leu Asn Gln Arg 245 250 255
Ala Ile Arg Ala Asp Thr Ser Tyr Cys Ile Arg Ile Thr Trp Ser Trp 260 265 270
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Thr Ser 290
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Thr Val Glu Gln Asp Ser Gly Gln Leu Ile Ala Asn Pro Lys Ala Pro 50 55 60

- Leu Gln Val Ala Asn Asp Lys Leu Glu Leu Ser Tyr Ala Tyr Pro Phe 65 70 75 80
- Glu Thr Ser Ala Asn Lys Leu Ser Leu Lys Val Gly Gln Gly Leu Lys 85 90 95
- Val Leu Asp Glu Lys Asp Ser Gly Gly Leu Gln Asn Leu Leu Gly Lys
  100 105 110
- Leu Val Val Leu Thr Gly Lys Gly Ile Gly Val Glu Glu Leu Lys Asn 115 120 125
- Pro Asp Asn Thr Asn Arg Gly Val Gly Ile Asn Val Arg Leu Gly Lys 130 135 140
- Asp Gly Gly Leu Ser Phe Asn Lys Asn Gly Glu Leu Val Ala Trp Asn 145 150 155 160
- Lys His Asn Asp Thr Gly Thr Leu Trp Thr Thr Pro Asp Pro Ser Pro 165 170 175
- Asn Cys Lys Ile Glu Glu Val Lys Asp Ser Lys Leu Thr Leu Val Leu 180 185 190
- Thr Lys Cys Gly Ser Gln Ile Leu Ala Thr Met Ala Phe Gln Val Val 195 200 205
- Lys Gly Thr Tyr Glu Asn Ile Ser Lys Asn Thr Ala Lys Asn Ser Phe 210 215 220
- Ser Ile Lys Leu Leu Phe Asp Asp Asn Gly Lys Leu Leu Glu Gly Ser 225 230 235 240
- Ser Leu Asp Lys Asp Tyr Trp Asn Phe Arg Ser Asp Asp Ser Ile Ile 245 250 255
- Pro Asn Gln Tyr Asp Asn Ala Val Pro Phe Met Pro Asn Leu Lys Ala 260 265 270
- Tyr Pro Lys Pro Ser Thr Val Leu Pro Ser Thr Asp Lys Asn Ser Asn 275 280 285
- Gly Lys Asn Thr Ile Val Ser Asn Leu Tyr Leu Glu Gly Lys Ala Tyr 290 295 300

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Gln Pro Val Ala Val Thr Ile Thr Phe Asn Lys Glu Ile Gly Cys Thr

Lys Arg Ala Arg Pro Ser Glu Asp Thr Phe Asn Pro Val Tyr Pro Tyr Gly Tyr Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Pro Pro Phe Val Ser Ser Asp Gly Phe Gln Asn Phe Pro Gly Val Leu Ser Leu Lys Leu Ala Asp Pro Ile Ala Ile Thr Asn Gly Asp Tyr Ser Leu Lys Val Gly Gly Gly Leu Thr Val Glu Lys Asp Ser Gly Asn Leu Lys Val Asn Pro Lys Ala Pro Leu Gln Val Thr Thr Asp Lys Gln Leu Glu Ile Ala Leu Ala Tyr Pro Phe Glu Val Ser Asn Gly Lys Leu Gly Ile Lys Ala Gly His Gly Leu Lys Val Ile Asp Lys Ile Ala Gly Leu Glu Gly Leu Ala Gly Thr Leu Val Val Leu Thr Gly Lys Gly Ile Gly Thr Glu Asn Leu Glu Asn Ser Asp Gly Ser Ser Arg Gly Val Gly Ile Asn Val Arg Leu Ala Lys Asp Gly Gly Leu Ser Phe Asp Lys Lys Gly Asp Leu Val Ala Trp Asn Lys His Asp Asp Arg Arg Thr Leu Trp Thr Thr Pro Asp Pro Ser Pro Asn Cys Thr Ile Asp Gln Glu Arg Asp Ser Lys Leu Thr Leu 

Val Leu Thr Lys Cys Gly Ser Gln Ile Leu Ala Asn Val Ser Leu Leu

Val Val Lys Gly Lys Phe Ser Asn Ile Asn Asn Asn Thr Asn Pro Thr

Asp Lys Lys Ile Thr Val Lys Leu Leu Phe Asn Glu Lys Gly Val Leu 260 265 270
Met Asp Ser Ser Thr Leu Lys Lys Glu Tyr Trp Asn Tyr Arg Asn Asp 275 280 285
Asn Ser Thr Val Ser Gln Ala Tyr Asp Asn Ala Val Pro Phe Met Pro 290 295 300
Asn Ile Lys Ala Tyr Pro Lys Pro Thr Thr Asp Thr Ser Ala Lys Pro 305 310 315 320
Glu Asp Lys Lys Ser Ala Ala Lys Arg Tyr Ile Val Ser Asn Val Tyr 325 330 335
Ile Gly Gly Leu Pro Asp Lys Thr Val Val Ile Thr Ile Lys Phe Asn 340 345 350
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Tyr Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Ile Pro Phe Val 35 40 45
Ser Ser Asp Gly Phe Gln Asn Phe Pro Pro Gly Val Leu Ser Leu Lys
50 55 60

Leu Ala Asp Pro Ile Thr Ile Ser Asn Gly Asp Val Ser Leu Lys Val Gly Gly Leu Thr Val Glu Gln Asp Ser Gly Asn Leu Ser Val Asn Pro Lys Ala Pro Leu Gln Val Gly Thr Asp Lys Leu Glu Leu Ala Leu Ala Pro Pro Phe Asn Val Lys Asp Asn Lys Leu Asp Leu Leu Val Gly Asp Gly Leu Lys Val Ile Asp Lys Ser Ile Ser Xaa Leu Pro Gly Leu Leu Asn Tyr Leu Val Val Leu Thr Gly Lys Gly Ile Gly Asn Glu Glu Leu Lys Leu Asp Asp Gly Ser Asn Lys Val Gly Leu Cys Val Arg Ile Gly Glu Gly Gly Leu Thr Phe Asp Asp Lys Gly Tyr Leu Val Ala Trp Asn Lys Lys His Asp Ile Arg Thr Leu Trp Thr Thr Leu Asp Pro Ser Pro Asn Cys Arg Ile Asp Val Asp Lys Asp Ser Lys Leu Thr Leu Val Leu Thr Lys Cys Gly Ser Gln Ile Leu Ala Asn Val Ser Leu Leu Val Val Lys Gly Arg Phe Gln Asn Leu Asn Tyr Lys Thr Asn Pro Asn Leu Pro Lys Thr Phe Thr Ile Lys Leu Leu Phe Asp Glu Asn Gly

Ile Leu Lys Asp Ser Ser Asn Leu Asp Lys Asn Tyr Trp Asn Tyr Arg

Asn Gly Asn Ser Ile Leu Ala Glu Gln Tyr Lys Asn Ala Val Gly Phe

Met Pro Asn Leu Ala Ala Tyr Pro Lys Ser Thr Thr Gln Ser Lys Leu Tyr Ala Arg Asn Thr Ile Phe Gly Asn Thr Tyr Leu Asp Ser Gln Ala Tyr Asn Pro Val Val Ile Lys Ile Thr Phe Asn Gln Glu Ala Asp Ser Ala Tyr Ser Ile Thr Leu Asn Tyr Ser Trp Gly Lys Asp Tyr Glu Asn Ile Pro Phe Asp Ser <210> 22 <211>334 <212> PRT <213> Human Adenovirus 27 Fiber Protein <400> 22 Ile Pro Phe Leu Thr Pro Pro Phe Val Ser Ser Asp Gly Phe Lys Asn Phe Pro Pro Gly Val Leu Ser Leu Lys Leu Ala Asp Pro Ile Thr Ile Thr Asn Gly Asp Val Ser Leu Lys Val Gly Gly Gly Leu Val Val Glu Lys Glu Ser Gly Lys Leu Ser Val Asp Pro Lys Thr Pro Leu Gln Val Ala Ser Asp Asn Lys Leu Glu Leu Ser Tyr Asn Ala Pro Phe Lys Val Glu Asn Asp Lys Leu Ser Leu Asp Val Gly His Gly Leu Lys Val Ile Gly Asn Glu Val Ser Ser Leu Pro Gly Leu Ile Asn Lys Leu Val Val Leu Thr Gly Lys Gly Ile Gly Thr Glu Leu Lys Glu Gln Asn Ser 

Asp Lys Ile Ile Gly Val Gly Ile Asn Val Arg Ala Arg Gly Gly Leu Ser Phe Asp Asn Asp Gly Tyr Leu Val Ala Trp Asn Pro Lys Tyr Asp Thr Arg Thr Leu Trp Thr Thr Pro Asp Thr Ser Pro Asn Cys Lys Met Leu Thr Lys Lys Asp Ser Lys Leu Thr Leu Thr Leu Thr Lys Cys Gly Ser Gln Ile Leu Gly Asn Val Ser Leu Leu Ala Val Ser Gly Lys Tyr Leu Asn Met Thr Lys Asp Glu Thr Gly Val Lys Ile Ile Leu Leu Phe Asp Arg Asn Gly Val Leu Met Gln Glu Ser Ser Leu Asp Lys Glu Tyr Trp Met Tyr Arg Asn Asp Asn Asn Val Ile Gly Thr Pro Tyr Glu Asn Ala Val Gly Phe Met Pro Asn Leu Val Ala Tyr Pro Lys Pro Thr Ser Ala Asp Ala Lys Asn Tyr Ser Arg Ser Lys Ile Ile Ser Asn Tyr Leu Lys Gly Leu Ile Tyr Gln Pro Val Ile Ile Ile Ala Ser Phe Asn Gln Glu Thr Thr Asn Gly Cys Val Tyr Ser Ile Ser Phe Asp Phe Thr Cys Ser Lys Asp Tyr Thr Gly Gln Gln Phe Asp Val Thr Ser Phe 

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Gly Tyr Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Pro Pro Phe Val 35 40 45
Ser Ser Asp Gly Phe Gln Asn Phe Pro Pro Gly Val Leu Ser Leu Lys 50 55 60
T ALA D TIME TI ALA CLA TILO TOTAL
Leu Ala Asp Pro Ile Thr Ile Ala Asn Gly Asp Val Ser Leu Lys Leu 65 70 75 80
Cly Cly Cly Cly I as The Wal Cly I are Cly Car Cly A - I - The Wal A
Gly Gly Leu Thr Val Glu Lys Glu Ser Gly Asn Leu Thr Val Asn 85 90 95
Des Less All, D. L. Cl. W. LAL C. Cl. Cl. L. Cl. V. LAL C.
Pro Lys Ala Pro Leu Gln Val Ala Ser Gly Gln Leu Glu Leu Ala Tyr 100 105 110
TC. D. DI A. VIII A. A. M. V. TILV. V. A. S.
Tyr Ser Pro Phe Asp Val Lys Asn Asn Met Leu Thr Leu Lys Ala Gly 115 120 125
It's Challen Ale Val Val That a A. A. Till A. I. Ch. D. I.
His Gly Leu Ala Val Val Thr Lys Asp Asn Thr Asp Leu Gln Pro Leu 130 135 140
M.C. W. I. W. I. W. I. W. C. Y. C. W. C. W.
Met Gly Thr Leu Val Val Leu Thr Gly Lys Gly Ile Gly Thr Gly Thr  145 150 155 160
143 130 133 100
Ser Ala His Gly Gly Thr Ile Asp Val Arg Ile Gly Lys Asn Gly Ser 165 170 175
Leu Ala Phe Asp Lys Asn Gly Asp Leu Val Ala Trp Asp Lys Glu Asn 180 185 190
Asp Arg Arg Thr Leu Trp Thr Thr Pro Asp Thr Ser Pro Asn Cys Lys 195 200 205
Met Ser Glu Val Lys Asp Ser Lys Leu Thr Leu Ile Leu Thr Lys Cys 210 215 220
Gly Ser Gln Ile Leu Gly Ser Val Ser Leu Leu Ala Val Lys Gly Glu 225 230 235 240

245 250 255
Leu Phe Asp Ala Asn Gly Val Leu Leu Glu Gly Ser Ser Leu Asp Lys 260 265 270
Glu Tyr Trp Asn Phe Arg Asn Asn Asp Ser Thr Val Ser Gly Lys Tyr 275 280 285
Glu Asn Ala Val Pro Phe Met Pro Asn Ile Thr Ala Tyr Lys Pro Val 290 295 300
Asn Ser Lys Ser Tyr Ala Arg Ser His Ile Phe Gly Asn Val Tyr Ile 305 310 315 320
Asp Ala Lys Pro Tyr Asn Pro Val Val Ile Lys Ile Ser Phe Asn Gln 325 330 335
Glu Thr Gln Asn Asn Cys Val Tyr Ser Ile Ser Phe Asp Tyr Thr Cys 340 345 350
Ser Lys Glu Tyr Thr Gly Met Gln Phe Asp Val Thr Ser Phe Thr Phe 355 360 365
Ser Tyr Ile Ala Gln Glu 370
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Ala Ile Thr Asn Gly Asp Val Ser Leu Lys Val Gly Gly Gly Leu Thr 35 40 45
Val Glu Gln Asp Ser Gly Asn Leu Ser Val Asn Pro Lys Ala Pro Leu 50 55 60

Gln Val Gly Thr Asp Lys Lys Leu Glu Leu Ala Leu Ala Pro Pro Phe 65 70 75 80
Asp Val Arg Asp Asn Lys Leu Ala Ile Leu Val Gly Asp Gly Leu Lys 85 90 95
Val Ile Asp Arg Ser Ile Ser Asp Leu Pro Gly Leu Leu Asn Tyr Leu 100 105 110
Val Val Leu Thr Gly Lys Gly Ile Gly Asn Glu Glu Leu Lys Asn Asp 115 120 125
Asp Gly Ser Asn Lys Gly Val Gly Leu Cys Val Arg Ile Gly Glu Gly 130 135 140
Gly Gly Leu Thr Phe Asp Asp Lys Gly Tyr Leu Val Ala Trp Asn Asr 145 150 155 160
Lys His Asp Ile Arg Thr Leu Trp Thr Thr Leu Asp Pro Ser Pro Asn 165 170 175
Cys Lys Ile Asp Ile Glu Lys Asp Ser Lys Leu Thr Leu Val Leu Thr 180 185 190
Lys Cys Gly Ser Gln Ile Leu Ala Asn Val Ser Leu Ile Ile Val Asn 195 200 205
Gly Lys Phe Lys Ile Leu Asn Asn Lys Thr Asp Pro Ser Leu Pro Lys 210 215 220
Ser Phe Asn Ile Lys Leu Leu Phe Asp Gln Asn Gly Val Leu Leu Glu 225 230 235 240
Asn Ser Asn Ile Glu Lys Gln Tyr Leu Asn Phe Arg Ser Gly Asp Ser 245 250 255
Ile Leu Pro Glu Pro Tyr Lys Asn Ala Ile Gly Phe Met Pro Asn Leu 260 265 270
Leu Ala Tyr Ala Lys Ala Thr Thr Asp Gln Ser Lys Ile Tyr Ala Arg 275 280 285
Asn Thr Thr Tyr Gly Asn Ile Tyr Leu Asp Asn Gln Pro Tyr Asn Pro 290 295 300

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Val Val Ile Lys Ile Thr Phe Asn Asn Glu Ala Asp Ser Ala Tyr Ser

Glu Leu Lys Asn Asp Asp Gly Ser Asn Lys Gly Val Gly Leu Cys Val 165 170 175
Arg Ile Gly Glu Gly Gly Leu Thr Xaa Asp Asp Lys Gly Tyr Leu 180 185 190
Val Ala Trp Asn Asn Lys His Asp Ile Arg Thr Leu Trp Thr Thr Leu 195 200 205
Asp Pro Ser Pro Asn Cys Lys Ile Asp Glu Lys Asp Ser Lys Leu Thr 210 215 220
Leu Val Leu Thr Lys Cys Gly Ser Gln Ile Leu Ala Asn Val Ser Leu 225 230 235 240
Ile Ile Val Asn Gly Lys Phe Lys Ile Leu Asn Asn Lys Thr Asp Pro 245 250 255
Ser Leu Pro Lys Ser Phe Asn Ile Lys Leu Leu Phe Asp Gln Asn Gly 260 265 270
Val Leu Leu Glu Asn Ser Asn Ile Glu Lys Gln Tyr Leu Asn Phe Arg 275 280 285
Ser Gly Asp Ser Ile Leu Pro Glu Pro Tyr Lys Asn Ala Ile Gly Phe 290 295 300
Met Pro Asn Leu Leu Ala Tyr Ala Lys Ala Thr Thr Asp Gln Ser Lys 305 310 315 320
Thr Tyr Ala Arg Asn Thr Ile Tyr Gly Asn Ile Tyr Leu Asp Asn Gln 325 330 335
Pro Tyr Asn Pro Val Val Ile Lys Ile Thr Phe Asn Asn Glu Ala Asp 340 345 350
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Asn Ile Pro Phe Asp Ser Thr Ser Phe Thr Phe Ser Tyr Ile Ala Gln 370 375 380
Glu

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                                 30
       20
Tyr Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Pro Pro Phe Val Ser
     35
                  40
                               45
Ser Asp Gly Phe Gln Asn Phe Pro Pro Gly Val Leu Ser Leu Lys Leu
   50
               55
                            60
Ala Asp Pro Ile Thr Ile Ala Asn Gly Asn Val Ser Leu Lys Val Gly
65
             70
                          75
                                       80
Gly Gly Leu Thr Leu Glu Gln Asp Ser Gly Lys Leu Ile Val Asn Pro
         85
                      90
                                   95
Lys Ala Pro Leu Gln Val Ala Asn Asp Lys Leu Glu Leu Ser Tyr Ala
       100
                    105
                                 110
Asp Pro Phe Glu Thr Ser Ala Asn Lys Leu Ser Leu Lys Val Gly His
    115
                  120
                               125
Gly Leu Lys Val Leu Asp Glu Lys Asn Ala Gly Gly Leu Lys Asp Leu
  130
               135
                             140
lle Gly Thr Leu Val Val Leu Thr Asp Lys Gly lle Gly Val Glu Glu
145
             150
                          155
                                        160
Leu Lys Asn Ala Asp Asn Thr Asn Arg Gly Val Gly Ile Asn Val Arg
         165
                      170
                                    175
Leu Gly Lys Asp Gly Gly Leu Ser Phe Asp Lys Lys Gly Asp Leu Val
                    185
      180
                                 190
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Ala Trp Asn Lys His Asp Asp Arg Arg Thr Leu Trp Thr Thr Pro Asp

210	215	220	i Glu Arg Asp Ser L	ys Leu Thr
Leu Val Le 225	u Thr Lys Cys 230	Gly Ser Gli 235	n Ile Leu Ala Asn Va 240	al Ser Leu
		Phe Ser Asi 250	n Ile Asn Asn Asn T 255	hr Asn Pro
Thr Asp Ly 260	s Lys Ile Thr 265	-	Leu Phe Asn Glu Ly	ys Gly Val
Leu Met As 275	sp Ser Ser Ser 280	Leu Lys Lys 285	Glu Tyr Trp Asn T	yr Arg Asn
Asp Asn Se 290	r Thr Ser Gln 295	Ala Tyr Asp 300	o Asn Ala Val Pro P	he Met Pro
Asn Ile Lys 305	Ala Tyr Pro I 310	Lys Pro Thr 7	Γhr Asp Thr Ser Ala 320	Lys Pro
Glu Asp Ly		Ala Lys Arg 30	Tyr Ile Val Ser Asr 335	ı Val Tyr
Ile Gly Gly	Leu Pro Asp I 345		Val Ile Thr Ile Lys I 50	eu Asn
Ala Glu Thr 355	Glu Ser Ala 7 360	Гуг Ser Met 365	Thr Phe Glu Phe Th	r Trp Ala
Lys Thr Phe 370	Glu Asn Leu 375	Gln Phe As 380	p Ser Ser Phe Tl	nr Phe Ser
Tyr Ile Ala ( 385	Gln Glu			
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_	Cys Pro Ser A		le Phe Met Leu Leu 15	Gin Met

Y ...

Lys Arg Ala Arg Pro Ser Glu Asp Thr Phe Asn Pro Val Tyr Pro Tyr 20 25 30
Gly Tyr Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Pro Pro Phe Val 35 40 45
Ser Ser Asp Gly Phe Lys Asn Phe Pro Pro Gly Val Leu Ser Leu Lys 50 55 60
Leu Ala Asp Pro Ile Thr Ile Thr Asn Gly Asp Val Ser Leu Lys Val 65 70 75 80
Gly Gly Leu Thr Leu Gln Glu Gly Ser Leu Thr Val Asn Pro Lys 85 90 95
Ala Pro Leu Gln Leu Ala Asn Asp Lys Lys Leu Glu Leu Val Tyr Asp 100 105 110
Asp Pro Phe Glu Val Ser Thr Asn Lys Leu Ser Leu Lys Val Gly His 115 120 125
Gly Leu Lys Val Leu Asp Asp Lys Ser Ala Gly Gly Leu Gln Asp Leu 130 135 140
Ile Gly Lys Leu Val Val Leu Thr Gly Lys Gly Ile Gly Ile Glu Asn 145 150 155 160
Leu Gln Asn Asp Asp Gly Ser Ser Arg Gly Val Gly Ile Asn Val Arg 165 170 175
Leu Gly Thr Asp Gly Gly Leu Ser Phe Asp Arg Lys Gly Glu Leu Val 180 185 190
Ala Trp Asn Arg Lys Asp Asp Arg Arg Thr Leu Trp Thr Thr Pro Asp 195 200 205
Pro Ser Pro Asn Cys Lys Ala Glu Thr Glu Lys Asp Ser Lys Leu Thr 210 215 220
Leu Val Leu Thr Lys Cys Gly Ser Gln Ile Leu Ala Thr Val Ser Ile 225 230 235 240
Ile Val Leu Lys Gly Lys Tyr Glu Phe Val Lys Lys Glu Thr Glu Pro 245 250 255

Lys Ser Phe A 260	sp Val Lys L 265		ne Asp Ser Lys Gly V 270	al Leu Leu
Pro Thr Ser A 275	sn Leu Ser Ly 280	ys Glu Tyr 285	r Trp Asn Tyr Arg Se 5	r Tyr Asp
Asn Asn Ile G 290	ly Thr Pro Ty 295	r Glu Ası 300	n Ala Val Pro Phe Me	et Pro Asn
-	yr Pro Lys Pr 310	o Thr Lys 315	s Thr Ala Ser Asp Lys 320	s Ala Glu
Asn Lys Ile Se 325	er Ser Ala Lys 330	_	Ile Val Ser Asn Phe	Tyr Phe
Gly Gly Gln A 340	da Tyr Gln Pi 345	-	r Ile Ile Ile Lys Phe A 350	sn Glu
Glu Ile Asp G	lu Thr Cys Al 360	la Tyr Ser 365	The Thr Phe Asn Phe	Gly Trp
Gly Lys Val T 370	yr Asp Asn P 375	Pro Phe Pr 380	o Phe Asp Thr Thr So	er Phe Thr
Phe Ser Tyr Ile 385	e Ala Gln Glu 390	1		
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Glu Asp Glu S 35	Ser Thr Ser G 40	ln His Pro 45	o Phe Ile Asn Pro Gly	Phe Ile
Ser Pro Asn G	ly Phe Thr G	ln Ser Pro	Asp Gly Val Leu Th	ır Leu Lys

Cys Leu Thr Pro Leu Thr Thr Gly Gly Ser Leu Gln Leu Lys Val 65 70 75 80
Gly Gly Leu Thr Val Asp Asp Thr Asp Gly Thr Leu Gln Lys Asn 85 90 95
Ile Arg Ala Thr Thr Pro Ile Thr Lys Asn Asn His Ser Val Glu Leu 100 105 110
Thr Ile Gly Asn Gly Leu Glu Thr Gln His Asn Lys Leu Cys Ala Lys 115 120 125
Leu Gly Asn Gly Asn Leu Lys Phe Asn Asn Gly Asp Ile Cys Ile Lys 130 135 140
Asp Ser Ile Asn Thr Leu Trp Thr Gly Ile Asn Pro Pro Asn Cys Gln 145 150 155 160
Ile Val Glu Asn Thr Asn Thr Asn Asp Gly Lys Leu Thr Leu Val Leu 165 170 175
Val Lys Asn Gly Gly Leu Val Asn Gly Tyr Val Ser Leu Val Gly Val 180 185 190
Ser Asp Thr Val Asn Gln Met Phe Thr Gln Lys Thr Ala Asn Ile Gln 195 200 205
Leu Arg Leu Tyr Phe Asp Ser Ser Gly Asn Leu Leu Thr Asp Glu Ser 210 215 220
Asp Leu Lys Ile Pro Leu Lys Asn Lys Ser Ser Thr Ala Thr Ser Glu 225 230 235 240
Thr Val Ala Ser Ser Lys Ala Phe Met Pro Ser Thr Thr Ala Tyr Pro 245 250 255
Phe Asn Thr Thr Arg Asp Ser Glu Asn Tyr Ile His Gly Ile Cys 260 265 270
Tyr Tyr Met Thr Ser Tyr Asp Arg Ser Leu Phe Pro Leu Asn Ile Ser 275 280 285
Ile Met Leu Asn Ser Arg Met Ile Ser Ser Asn Val Ala Tyr Ala Ile 290 295 300

Met Thr Leu Thr Thr Ser Pro Phe Phe Phe Ser Tyr Ile Ile Glu Asp Asp Asn <210> 29 <211>338 <212> PRT <213> Human Adenovirus 35 Fiber Protein <400> 29 Ser Cys Ser Cys Pro Ser Ala Pro Thr Ile Phe Met Leu Leu Gln Met Lys Arg Ala Arg Pro Ser Glu Asp Thr Phe Asn Pro Val Tyr Pro Tyr Glu Asp Glu Ser Thr Ser Gln His Pro Phe Ile Asn Pro Gly Phe Ile Ser Pro Asn Gly Phe Thr Gln Ser Pro Asp Gly Val Leu Thr Leu Lys Cys Leu Thr Pro Leu Thr Thr Gly Gly Ser Leu Gln Leu Lys Val Gly Gly Leu Thr Val Asp Asp Thr Asp Gly Thr Leu Gln Glu Asn Ile Arg Ala Thr Ala Pro Ile Thr Lys Asn Asn His Ser Val Glu Leu Ser Ile Gly Asn Gly Leu Glu Thr Gln Asn Asn Lys Leu Cys Ala Lys Leu Gly Asn Gly Leu Lys Phe Asn Asn Gly Asp Ile Cys Ile Lys Asp Ser Ile Asn Thr Leu Trp Thr Gly Ile Asn Pro Pro Pro Asn Cys Gln 

Gln Phe Glu Trp Asn Leu Asn Ala Ser Glu Ser Pro Glu Lys Gln His

Ile Val Gl	u Asn Thr 165	Asn Thr Ası 170	n Asp Gly Lys I 175	Leu Thr Leu Val Leu
Val Lys A		Leu Val As 185	n Gly Tyr Val 190	Ser Leu Val Gly Val
Ser Asp T 195		Gln Met Ph 00	ne Thr Gln Lys 205	Thr Ala Asn Ile Gln
Leu Arg L 210	eu Tyr Phe 215	-	r Gly Asn Leu 20	Leu Thr Glu Glu Ser
Asp Leu I 225	Lys Ile Pro	Leu Lys Asr 235	Lys Ser Ser Tl 240	nr Ala Thr Ser Glu
	la Ser Ser l 245	Lys Ala Phe 250	Met Pro Ser TI 255	nr Thr Ala Tyr Pro
Phe Asn T		Arg Asp Se 265	er Glu Asn Tyr 270	lle His Gly Ile Cys
Tyr Tyr M 275		Tyr Asp Arg 80	g Ser Leu Phe F 285	ro Leu Asn Ile Ser
Ile Met Le 290	eu Asn Ser 295	Arg Met Ile 30		ıl Ala Tyr Ala Ile
Gln Phe G	du Trp Asr 310	Leu Asn Al	la Ser Glu Ser I 320	Pro Glu Ser Asn Ile
	eu Thr Thr 325	Ser Pro Phe	Phe Phe Ser T	yr Ile Thr Glu Asp
Asp Asn				
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<213> Hu	man Adeno	ovirus 36 Fib	er Protein	
<400> 30				
Ser Cys Se	er Cys Pro	Ser Ala Pro	Thr Ile Phe Me	t Leu Leu Gln Met
1	5	10	15	

2.4

Υ.

Lys Arg Ala Arg Pro Ser Glu Asp Thr Phe Asn Pro Val Tyr Pro Tyr Gly Tyr Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Pro Pro Phe Val Ser Ser Asp Gly Phe Lys Asn Phe Pro Pro Gly Val Leu Ser Leu Lys Leu Ala Asp Pro Ile Ala Ile Val Asn Gly Asp Val Ser Leu Lys Val Gly Gly Leu Thr Val Glu Gln Asp Ser Gly Lys Leu Lys Val Asn Pro Lys Ile Pro Leu Gln Val Val Asn Lys Gln Leu Glu Leu Ala Thr Asp Lys Pro Phe Lys Ile Glu Asn Asn Lys Leu Ala Leu Asp Val Gly His Gly Leu Lys Val Ile Asp Lys Thr Ile Ser Asp Leu Gln Gly Leu Val Gly Lys Leu Val Val Leu Thr Gly Val Gly Ile Gly Thr Glu Thr Leu Lys Asp Lys Asn Asp Lys Val Ile Gly Ser Ala Val Asn Val Arg Leu Gly Lys Asp Gly Gly Leu Asp Phe Asn Lys Lys Gly Asp Leu Val Ala Trp Asn Arg Tyr Asp Asp Arg Arg Thr Leu Trp Thr Thr Pro Asp Pro Ser Pro Asn Cys Lys Val Tyr Glu Ala Lys Ser Lys Leu Thr Leu Val Leu Thr Lys Cys Gly Ser Gln Ile Leu Ala Ser Val Ala Leu Leu 

Ile Val Lys Gly Lys Tyr Gln Thr Ile Ser Glu Ser Thr Ile Pro Lys

Asp Gln Arg Asn Phe Ser Val Lys Leu Met Phe Asp Glu Lys Gly Lys 260 265 270
Leu Leu Asp Lys Ser Ser Leu Asp Lys Glu Tyr Trp Asn Phe Arg Ser 275 280 285
Asn Asp Ser Val Val Gly Thr Ala Tyr Asp Asn Ala Val Pro Phe Met 290 295 300
Pro Asn Leu Lys Ala Tyr Pro Lys Asn Thr Thr Thr Ser Ser Thr Asn 305 310 315 320
Pro Asp Asp Lys Ile Ser Ala Gly Lys Lys Asn Ile Val Ser Asn Val 325 330 335
Tyr Leu Glu Gly Arg Val Tyr Gln Pro Val Ala Leu Thr Val Lys Phe 340 345 350
Asn Ser Glu Asn Asp Cys Ala Tyr Ser Ile Thr Phe Asp Phe Val Trp 355 360 365
Ser Lys Thr Tyr Glu Ser Pro Val Ala Phe Asp Ser Ser Ser Phe Thr 370 375 380
Phe Ser Tyr Ile Ala Gln Glu 385 390
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Gly Tyr Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Pro Pro Phe Val 35 40 45
Ser Ser Asp Gly Phe Lys Asn Phe Pro Pro Gly Val Leu Ser Leu Lys

Leu Ala Asp Pro Ile Thr Ile Thr Asn Gly Asp Val Ser Leu Lys Val 65 70 75 80
Gly Gly Leu Thr Leu Gln Asp Gly Ser Leu Thr Val Asn Pro Lys 85 90 95
Ala Pro Leu Gln Val Asn Thr Asp Lys Lys Leu Glu Leu Ala Tyr Asp 100 105 110
Asn Pro Phe Glu Ser Ser Ala Asn Lys Leu Ser Leu Val Gly His Gly 115 120 125
Leu Lys Val Leu Asp Glu Lys Ser Ala Ala Gly Leu Lys Asp Leu Ile 130 135 140
Gly Lys Leu Val Val Leu Thr Gly Lys Gly Ile Gly Thr Glu Asn Leu 145 150 155 160
Glu Asn Thr Asp Gly Ser Ser Arg Gly Ile Gly Ile Asn Val Arg Ala 165 170 175
Arg Glu Gly Leu Thr Phe Asp Asn Asp Gly Tyr Leu Val Ala Trp Asn 180 185 190
Pro Lys Tyr Asp Leu Arg Thr Leu Trp Thr Thr Pro Asp Thr Ser Pro 195 200 205
Asn Cys Thr Ile Ala Gln Asp Lys Asp Ser Lys Leu Thr Leu Val Leu 210 215 220
Thr Lys Cys Gly Ser Gln IIe Leu Ala Asn Val Ser Leu IIe Val Val 225 230 235 240
Ala Gly Lys Tyr His Ile Ile Asn Asn Lys Thr Asn Pro Lys Ile Lys 245 250 255
Ser Phe Thr Ile Lys Leu Leu Phe Asn Lys Phe Asn Gly Val Leu Leu 260 265 270
Asp Asn Ser Asn Leu Gly Lys Ala Tyr Trp Asn Phe Arg Ser Gly Asn 275 280 285
Ser Asn Val Ser Thr Ala Tyr Glu Lys Ala Ile Gly Phe Met Pro Asn

Leu Val Ala Val Ser Lys Pro Ser Asn Ser Lys Lys Tyr Ala Arg Asp 305 310 315 320
Ile Val Tyr Gly Asn Ile Thr Tyr Leu Gly Gly Lys Pro Asp Gln Pro 325 330 335
Gly Val Ile Lys Thr Thr Phe Asn Gln Glu Thr Gly Cys Glu Tyr Ser 340 345 350
Ile Thr Phe Asn Phe Ser Trp Ser Lys Thr Tyr Glu Asn Val Glu Phe 355 360 365
Glu Thr Thr Ser Phe Thr Phe Ser Tyr Ile Ala Gln Glu 370 375 380
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Gly Tyr Ala Arg Asn Gln Asn Ile Pro Phe Xaa Thr Pro Pro Phe Val 35 40 45
Xaa Ser Asp Gly Phe Gln Asn Phe Pro Pro Gly Val Leu Ser Leu Lys 50 55 60
Leu Ala Asp Pro Ile Thr Ile Ala Asn Gly Asn Val Ser Leu Lys Val 65 70 75 80
Gly Gly Leu Thr Leu Glu Gln Asp Ser Gly Lys Leu Ile Val Asn 85 90 95
Xaa Lys Ala Pro Leu Gln Val Ala Asn Asp Lys Leu Glu Leu Ser Tyr 100 105 110
Ala Asp Pro Phe Glu Thr Ser Ala Asn Lys Leu Ser Leu Lys Val Gly 115 120 125

His Gly Leu Lys Val Leu Asp Glu Lys Asn Ala Gly Gly Leu Lys Asp 130 135 140
Leu Ile Gly Thr Leu Val Val Leu Thr Gly Lys Gly Ile Gly Val Glu 145 150 155 160
Glu Leu Lys Asn Ala Asp Asn Thr Asn Arg Gly Val Gly Ile Asn Val 165 170 175
Arg Leu Gly Lys Asp Gly Gly Leu Ser Phe Asp Lys Lys Gly Asp Xaa 180 185 190
Val Ala Trp Asn Lys His Asp Asp Arg Arg Thr Leu Trp Thr Thr Pro 195 200 205
Asp Pro Ser Pro Asn Cys Thr Ile Asp Glu Glu Arg Asp Ser Lys Leu 210 215 220
Thr Leu Val Leu Thr Lys Cys Gly Ser Gln Ile Leu Ala Asn Val Ser 225 230 235 240
Leu Leu Val Val Lys Gly Lys Phe Ser Asn Ile Asn Asn Asn Thr Asn 245 250 255
Pro Thr Asp Lys Lys Ile Thr Val Lys Leu Leu Phe Asn Glu Lys Gly 260 265 270
Val Leu Met Asp Ser Ser Ser Leu Lys Lys Glu Tyr Trp Asn Tyr Arg 275 280 285
Asn Asp Asn Ser Thr Val Ser Gln Ala Tyr Asp Asn Ala Val Pro Phe 290 295 300
Met Pro Asn Ile Lys Ala Tyr Pro Lys Pro Thr Thr Asp Thr Ser Ala 305 310 315 320
Lys Pro Glu Asp Lys Lys Ser Ala Ala Lys Arg Tyr Thr Val Ser Asn 325 330 335
Val Tyr Ile Gly Gly Leu Pro Asp Lys Thr Val Val Ile Thr Ile Lys 340 345 350
Leu Asn Ala Glu Thr Glu Ser Ala Tyr Ser Met Thr Phe Glu Phe Thr 355 360 365

Phe Ser Tyr Ile Ala Gln Glu <210> 33 <211>338 <212> PRT <213> Human Adenovirus 39 Fiber Protein <400> 33 Ile Arg Ile Ser Pro Ser Ser Leu Pro Pro Leu Ser Pro Pro Met Asp Ser Lys Thr Ser Pro Leu Gly Cys Tyr His Ser Asn Trp Leu Thr Gln Ser Pro Ser Pro Met Gly Met Ser His Arg Trp Glu Gly Gly Ser Pro Trp Gln Glu Gly Thr Gly Asp Leu Lys Val Asn Ala Lys Ser Pro Leu Gln Val Ala Thr Asn Lys Gln Leu Glu Ile Ala Leu Ala Lys Pro Phe Glu Glu Lys Asp Gly Lys Leu Ala Leu Lys Ile Gly His Gly Leu Ala Val Val Asp Glu Asn His Thr His Leu Gln Ser Leu Ile Gly Thr Leu Val Ile Leu Thr Gly Lys Gly Ile Gly Thr Gly Arg Ala Glu Ser Gly Gly Thr Ile Asp Val Arg Leu Gly Ser Gly Gly Leu Ser Phe Asp Lys Asp Gly Asn Leu Val Ala Trp Asn Lys Asp Asp Asp Arg Arg Thr Leu Trp Thr Thr Pro Asp Pro Ser Pro Asn Cys Lys Ile Asp Gln Asp 

Trp Ala Lys Thr Phe Glu Asn Leu Gln Phe Asp Ser Ser Ser Phe Thr

Lys Asp Ser Lys Leu Thr Phe Val Leu Thr Lys Cys Gly Ser Gln Ile 180 185 190
Leu Ala Asn Met Ser Leu Leu Val Val Lys Gly Lys Phe Ser Met Ile 195 200 205
Asn Asn Lys Val Asn Gly Thr Asp Asp Tyr Lys Lys Phe Thr Ile Lys 210 215 220
Leu Leu Phe Asp Glu Lys Gly Val Leu Leu Lys Asp Ser Ser Leu Asp 225 230 235 240
Lys Glu Tyr Trp Asn Tyr Arg Ser Asn Asn Asn Asn Val Gly Ser Ala 245 250 255
Tyr Glu Glu Ala Val Gly Phe Met Pro Ser Thr Thr Ala Tyr Pro Lys 260 265 270
Pro Pro Thr Pro Pro Thr Asn Pro Thr Thr Pro Leu Glu Lys Ser Gln 275 280 285
Ala Lys Asn Lys Tyr Val Ser Asn Val Tyr Leu Gly Gly Gln Ala Gly 290 295 300
Asn Pro Val Ala Thr Thr Val Ser Phe Asn Lys Glu Thr Gly Cys Thr 305 310 315 320
Tyr Ser Ile Thr Phe Asp Phe Ala Trp Asn Lys Thr Tyr Glu Asn Val 325 330 335
Gln Cys
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<400> 34 Ser Cys Ser Cys Pro Ser Ala Pro Thr Ile Phe Met Leu Leu Gln Met
1 5 10 15
Lys Arg Ala Arg Pro Ser Glu Asp Thr Phe Asn Pro Val Tyr Pro Tyr 20 25 30

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Gly Tyr Ala Arg Asr 35 4		o Phe Leu Thr I	Pro Pro Phe Val
Ser Ser Asp Gly Phe 50 55	Lys Asn Phe P	ro Pro Gly Val I	Leu Ser Leu Lys
Leu Ala Asn Pro Ile 65 70	Ala Ile Thr Asn 75	Gly Asp Val Se 80	er Leu Lys Val
Gly Gly Gly Leu Thr 85	Leu Gln Asp C 90	Gly Thr Gly Lys 95	Leu Thr Ile Asp
Thr Lys Thr Pro Leu 100	Gln Val Ala As 105	sn Asn Lys Leu 110	Glu Leu Ala Phe
Asp Ala Pro Leu Tyr 115 1:	Glu Lys Asn C 20 12	• •	Leu Lys Thr Gly
His Gly Leu Ala Val 130 135	Leu Thr Lys A	sp Ile Gly Ile Pr	o Glu Leu Ile
Gly Ser Leu Val Ile I 145 150	eu Thr Gly Lys. 155	s Gly Ile Gly Th 160	r Gly Thr Val
Ala Gly Gly Gly Thr 165	Ile Asp Val Ar	g Leu Gly Asp A	Asp Gly Gly Leu
Ser Phe Asp Lys Lys 180	-	'al Ala Trp Asn 190	Lys Lys Asn Asp
Arg Arg Thr Leu Trp	Thr Thr Pro A	_	Asn Cys Arg Val
Ser Glu Asp Lys Asp 210 215	Ser Lys Leu Ti 220	hr Leu Ile Leu T	hr Lys Cys Gly
Ser Gln Ile Leu Ala S 225 230	Ser Phe Ser Leu 235	Leu Val Val Xa 240	a Gly Thr Tyr
Thr Thr Val Asp Lys 245	Asn Thr Thr A 250	sn Lys Gln Phe 255	Ser Ile Lys Leu
Leu Phe Asp Ala Ası 260	• •	.ys Ser Glu Ser . 270	Asn Leu Ser Gln

Tyr Trp Asn Tyr Arg Ser Asp Asn Ser Val Val Ser Thr Pro Tyr Asp 275 280 285
Asn Ala Val Pro Phe Met Pro Asn Thr Ala Tyr Pro Lys Ile Ile Asn 290 295 300
Ser Thr Thr Asp Pro Glu Asn Lys Lys Ser Ala Lys Lys Thr Ile Val 305 310 315 320
Gly Asn Val Tyr Leu Glu Gly Asn Ala Gly Gln Pro Val Ala Val Ala 325 330 335
Ile Ser Phe Asn Lys Glu Thr Thr Ala Asp Tyr Ser Ile Thr Phe Asp 340 345 350
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Ile Thr Asn Gly Asp Val Ser Leu Lys Val Gly Gly Gly Leu Thr Val 35 40 45
Glu Lys Glu Ser Gly Asn Leu Thr Val Asn Pro Lys Ala Pro Leu Gln 50 55 60
Val Ala Lys Gly Gln Leu Glu Leu Ala Tyr Asp Ser Pro Phe Asp Val 65 70 75 80
Lys Asn Asn Met Leu Thr Leu Lys Ala Gly His Gly Leu Ala Val Val 85 90 95

Thr Lys Asp Asn Thr Asp Leu Gln Pro Leu Met Gly Thr Leu Val Val 100 105 110
Leu Thr Gly Lys Gly Ile Gly Thr Gly Thr Ser Ala His Gly Gly Thr 115 120 125
Ile Asp Val Arg Ile Gly Lys Asn Gly Ser Leu Ala Phe Asp Lys Asp 130 135 140
Gly Asp Leu Val Ala Trp Asp Lys Glu Asn Asp Arg Arg Thr Leu Trp 145 150 155 160
Thr Thr Pro Asp Thr Ser Pro Asn Cys Lys Met Ser Glu Ala Lys Asp 165 170 175
Ser Lys Leu Thr Leu Ile Leu Thr Lys Cys Gly Ser Gln Ile Leu Gly 180 185 190
Ser Val Ser Leu Leu Ala Val Lys Gly Glu Tyr Gln Asn Met Thr Ala 195 200 205
Asn Thr Lys Lys Asn Val Lys Ile Thr Leu Leu Phe Asp Ala Asn Gly 210 215 220
Val Leu Leu Ala Gly Ser Ser Xaa Xaa Lys Glu Tyr Trp Asn Phe Arg 225 230 235 240
Ser Asn Asp Ser Thr Val Ser Gly Asn Tyr Glu Asn Ala Val Gln Phe 245 250 255
Met Pro Asn Ile Thr Ala Tyr Lys Pro Thr Asn Ser Lys Ser Tyr Ala 260 265 270
Arg Ser Val Ile Phe Gly Asn Val Tyr Ile Asp Ala Lys Pro Tyr Asn 275 280 285
Pro Val Val Ile Lys Ile Ser Phe Asn Gln Glu Thr Gln Asn Asn Cys 290 295 300
Val Tyr Ser Ile Ser Phe Asp Tyr Thr Leu Ser Lys Asp Tyr Pro Asn 305 310 315 320
Met Gln Phe Asp Val Thr Leu Ser 325

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                                 30
       20
Ile Thr Asn Gly Asn Val Ser Leu Lys Val Gly Gly Leu Thr Leu
     35
                 40
                              45
Gln Glu Gly Thr Gly Asp Leu Lys Val Asn Ala Lys Ser Pro Leu Gln
                            60
  50
               55
Val Ala Thr Asn Lys Gln Leu Glu Ile Ala Leu Ala Lys Pro Phe Glu
65
             70
                         75
                                      80
Glu Lys Asp Gly Lys Leu Ala Leu Lys Ile Gly His Gly Leu Ala Val
         85
                      90
                                   95
Val Asp Glu Asn His Thr His Leu Gln Ser Leu Ile Gly Thr Leu Val
                    105
                                 110
      100
Ile Leu Thr Gly Lys Gly Ile Gly Thr Gly Ser Ala Glu Ser Gly Gly
    115
                 120
                               125
Thr Ile Asp Val Arg Leu Gly Ser Gly Gly Leu Ser Phe Asp Lys
  130
               135
                             140
Asp Gly Asn Leu Val Ala Trp Asn Lys Asp Asp Asp Arg Arg Thr Leu
                                        160
145
             150
                          155
Trp Thr Thr Pro Asp Pro Ser Pro Asn Cys Lys Ile Asp Gln Asp Lys
         165
                      170
                                    175
Asp Ser Lys Leu Thr Phe Val Leu Thr Lys Cys Gly Ser Gln Ile Leu
      180
                    185
                                 190
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Ala Asn Met Ser Leu Leu Val Val Lys Gly Lys Phe Ser Met Ile Asn

Asn Lys Val Asn Gly Thr Asp Asp Tyr Lys Lys Phe Thr Ile Lys Leu 210 215 220
Leu Phe Asp Glu Lys Gly Val Leu Leu Lys Asp Ser Ser Leu Asp Lys 225 230 235 240
Glu Tyr Trp Asn Tyr Arg Ser Asn Asn Asn Asn Val Gly Ser Ala Tyr 245 250 255
Glu Glu Ala Val Gly Phe Met Pro Ser Thr Thr Ala Tyr Pro Lys Pro 260 265 270
Pro Thr Pro Pro Thr Asn Pro Thr Thr Pro Leu Glu Lys Ser Gln Ala 275 280 285
Lys Asn Lys Tyr Val Ser Asn Val Tyr Leu Gly Gly Gln Ala Gly Asn 290 295 300
Pro Val Ala Thr Thr Val Ser Phe Asn Lys Glu Thr Gly Cys Thr Tyr 305 310 315 320
Ser Ile Thr Phe Asp Phe Ala Trp Asn Lys Thr Tyr Glu Asn Val Gln 325 330 335
Phe Asp Ser Ser Phe 340
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Ile Thr Asn Gly Asp Val Ser Leu Lys Val Gly Gly Gly Leu Thr Val 35 40 45
Glu Lys Asp Ser Gly Asn Leu Lys Val Asn Pro Lys Ala Pro Leu Gln 50 55 60

Val Thr Thr Asp Lys Gln Leu Glu Ile Ala Leu Ala Tyr Pro Phe Glu 65 70 75 80
Val Ser Asn Gly Lys Leu Gly Ile Lys Ala Gly His Gly Leu Lys Val 85 90 95
Ile Asp Lys Ile Ala Gly Leu Glu Gly Leu Ala Gly Thr Leu Val Val 100 105 110
Leu Thr Gly Lys Gly Ile Gly Thr Glu Asn Leu Glu Asn Ser Asp Gly 115 120 125
Ser Ser Arg Gly Val Gly Ile Asn Val Arg Leu Ala Lys Asp Gly Val 130 135 140
Leu Ala Phe Asp Lys Lys Gly Asp Leu Val Ala Trp Asn Lys His Asp 145 150 155 160
Asp Arg Arg Thr Leu Trp Thr Thr Pro Asp Pro Ser Pro Asn Cys Thr 165 170 175
Ile Asp Gln Glu Arg Asp Ser Lys Leu Thr Leu Val Leu Thr Lys Cys 180 185 190
Gly Ser Gln Ile Leu Ala Asn Val Ser Leu Leu Val Val Lys Gly Lys 195 200 205
Phe Ser Asn Ile Asn Asn Asn Ala Asn Pro Thr Asp Lys Lys Ile Thr 210 215 220
Val Lys Leu Leu Phe Asn Glu Lys Gly Val Leu Met Asp Ser Ser Thr 225 230 235 240
Leu Lys Lys Glu Tyr Trp Asn Tyr Arg Asn Asp Asn Ser Thr Val Ser 245 250 255
Gln Ala Tyr Asp Asn Ala Val Pro Phe Met Pro Asn Ile Lys Ala Tyr 260 265 270
Pro Lys Pro Ser Thr Asp Thr Ser Ala Lys Pro Glu Asp Lys Lys Ser 275 280 285
Ala Ala Lys Arg Tyr Ile Val Ser Asn Val Tyr Ile Gly Gly Leu Pro

Ala Tyr Ser Ile Thr Phe Glu Phe Thr Trp Ala Lys Thr Phe Glu Asp Val Gln Cys Asp Ser Ser Ser Phe Thr <210>38 <211>339 <212> PRT <213> Human Adenovirus 46 Fiber Protein <400>38 Asn Ile Pro Phe Leu Thr Pro Pro Phe Val Ser Ser Asp Gly Phe Lys Asn Phe Pro Pro Gly Val Leu Ser Leu Lys Leu Ala Asp Pro Ile Ala Ile Val Asn Gly Asp Val Ser Leu Lys Val Gly Gly Leu Thr Leu Gln Glu Gly Asn Leu Thr Val Asp Ala Lys Ala Pro Leu Gln Val Ala Asn Asp Lys Leu Glu Leu Ser Tyr Ala Asp Phe Phe Glu Val Lys Asp Thr Lys Leu Gln Leu Lys Val Gly His Gly Leu Lys Val Ile Asp Glu Lys Thr Ser Ser Gly Leu Gln Ser Leu Ile Gly Asn Leu Val Val Leu Thr Gly Lys Gly Ile Gly Thr Gln Glu Leu Lys Asp Lys Asp Glu Thr Lys Asn Ile Gly Val Gly Ile Asn Val Arg Ile Gly Lys Asn Glu Ser Leu Ala Phe Asp Lys Asp Gly Asn Leu Val Ala Trp Asp Asn Glu 

Asp Lys Thr Val Val Ile Thr Ile Lys Phe Asn Ala Glu Thr Glu Cys

Asn Asp Arg Arg Thr Leu Trp Thr Thr Pro Asp Thr Ser Ser Lys Phe 165 170 175
Val Lys Ile Ser Thr Glu Lys Asp Ser Lys Leu Thr Leu Val Leu Thr 180 185 190
Lys Cys Gly Ser Gln Ile Leu Ala Ser Val Ser Leu Leu Ala Val Ala 195 200 205
Gly Ser Tyr Leu Asn Met Thr Ala Ser Thr Gln Lys Ser Ile Lys Val 210 215 220
Ser Leu Met Phe Asp Ser Lys Gly Leu Leu Met Thr Thr Ser Ser Ile 225 230 235 240
Asp Lys Gly Tyr Trp Asn Tyr Arg Asn Lys Asn Ser Val Val Gly Thr 245 250 255
Ala Tyr Glu Asn Ala Ile Pro Phe Met Pro Asn Leu Val Ala Tyr Pro 260 265 270
Arg Pro Asn Thr Pro Asp Ser Lys Ile Tyr Ala Arg Ser Lys Ile Val 275 280 285
Gly Asn Val Tyr Leu Ala Gly Leu Ala Tyr Gln Pro Ile Val Ile Thr 290 295 300
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Ser Phe Thr
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Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Pro Pro Phe Val Ser Ser 35 40 45	
Asp Gly Phe Lys Asn Phe Pro Pro Gly Val Leu Ser Leu Lys Leu Als 50 55 60	a
Asp Pro Ile Thr Ile Thr Asn Gly Asp Val Ser Leu Lys Val Gly Gly 65 70 75 80	
Gly Leu Thr Leu Gln Glu Gly Thr Gly Asn Leu Thr Val Asn Ala Ly 85 90 95	/S
Ala Pro Leu Gln Val Ala Asp Asp Lys Lys Leu Glu Leu Ser Tyr Asj 100 105 110	p
Asn Pro Phe Glu Val Ser Ala Asn Lys Leu Ser Leu Lys Val Gly His 115 120 125	•
Gly Leu Lys Val Leu Asp Glu Lys Asn Ser Gly Gly Leu Gln Glu Le 130 135 140	u
Ile Gly Lys Leu Val Ile Leu Thr Gly Lys Gly Ile Gly Val Glu Glu 145 150 155 160	
Leu Lys Asn Ala Asp Asn Thr Asn Arg Gly Val Gly Ile Asn Val Arg 165 170 175	g
Leu Gly Lys Asp Gly Gly Leu Ser Phe Asp Lys Lys Gly Glu Leu Va 180 185 190	ıl
Ala Trp Asn Lys His Asn Asp Thr Arg Thr Leu Trp Thr Thr Pro Asj 195 200 205	p
Pro Ser Pro Asn Cys Lys Ile Glu Gln Asp Lys Asp Ser Lys Leu Thr 210 215 220	
Leu Val Leu Thr Lys Cys Gly Ser Gln Ile Leu Ala Thr Met Ala Phe 225 230 235 240	
Gln Val Val Lys Asp Thr Tyr Glu Asn Ile Ser Lys Asn Thr Ala Lys 245 250 255	

Lys Ser Phe Ser Ile Lys Leu Leu Phe Asp Asp Asn Gly Lys Leu Leu 260 265 270
Glu Gly Ser Ser Leu Asp Lys Asp Tyr Trp Asn Phe Arg Asn Asp Asp 275 280 285
Ser Ile Met Pro Ser Gln Tyr Asp Asn Ala Val Pro Phe Met Pro Asn 290 295 300
Leu Lys Ala Tyr Pro Asn Pro Lys Thr Ser Thr Val Leu Pro Ser Thr 305 310 315 320
Asp Lys Lys Ser Asn Gly Lys Asn Thr Ile Val Ser Asn Leu Tyr Leu 325 330 335
Glu Gly Lys Ala Tyr Gln Pro Val Ala Val Thr Ile Thr Phe Asn Lys 340 345 350
Glu Tyr Gly Cys Thr Tyr Ser Ile Thr Phe Glu Phe Gly Trp Ala Lys 355 360 365
Thr Tyr Asp Val Pro Ile Pro Phe Asp Ser Ser Ser Phe Thr Phe Ser 370 375 380
Tyr Ile Ala Gln Glu 385
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<400> 40 Ser Asp Ile Pro Phe Leu Thr Pro Pro Phe Val Ser Ser Asp Gly Phe 1 5 10 15
Gln Asn Phe Pro Pro Gly Val Leu Ser Leu Lys Leu Ala Asp Pro Ile 20 25 30
Thr Ile Thr Asn Gly Asn Val Ser Leu Lys Val Gly Gly Gly Leu Thr 35 40 45
Leu Gln Glu Gly Thr Gly Asp Leu Lys Val Asn Ala Lys Ser Pro Leu 50 55 60

Gln Val Ala Thr Asn l 65 70	Lys Gln Leu Glu 75	Ile Ala Leu Ala Lys Pro Phe 80
Glu Glu Lys Asp Gly 1 85	-	Lys Ile Gly His Glu Leu Ala 95
-	Leu Thr His Leu 105 11	Gln Ser Leu Ile Gly Thr Leu 0
Val Ile Leu Thr Gly Ly 115 120	•	nr Gly Arg Ala Glu Ser Gly
Gly Thr Ile Asp Val A 130 135	rg Leu Gly Ser ( 140	Gly Gly Gly Leu Ser Phe Asp
Lys Asp Gly Asn Leu 145 150	Val Ala Trp Asn 155	Lys Asp Asp Asp Arg Arg Thr 160
Leu Trp Thr Thr Pro A	_	Asn Cys Lys Ile Asp Gln Asp 175
	Thr Phe Val Leu 185 19	Thr Lys Cys Gly Ser Gln Ile
Leu Ala Asn Met Ser I 195 200		Lys Gly Lys Phe Ser Met Ile
Asn Asn Lys Val Asn 210 215	Gly Thr Asp As <sub>l</sub> 220	p Tyr Lys Lys Phe Thr Ile Lys
Leu Leu Phe Asp Glu l 225 230	Lys Gly Val Leu 235	Leu Lys Asp Ser Ser Leu Asp 240
Lys Glu Tyr Trp Asn T 245	•	Asn Asn Asn Val Gly Ser Ala 255
-	Gly Phe Met Pro 265 279	Ser Thr Thr Ala Tyr Pro Lys 0
Pro Pro Thr Pro Pro Th 275 280		hr Pro Leu Glu Lys Ser Gln
Ala Lys Asn Lys Tyr V 290 295	Val Ser Asn Val	Гуг Leu Gly Gly Gln Ala Gly

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Tyr Ser Ile Thr Phe Asp Phe Ala Trp Asn Lys Thr Tyr Lys Met Ala Phe Ile Pro Arg Phe Asn Phe <210>41 <211>393 <212> PRT <213> Human Adenovirus 49 Fiber Protein <400> 41 Ser Cys Ser Cys Pro Ser Ala Pro Thr Ile Phe Met Leu Gln Met Lys Arg Ala Arg Pro Ser Glu Asp Thr Phe Asn Pro Val Tyr Pro Tyr Gly Tyr Ala Arg Asn Gln Asn Ile Pro Phe Leu Thr Pro Pro Phe Val Ser Ser Asp Gly Phe Gln Asn Phe Pro Pro Gly Val Leu Ser Leu Lys Leu Ala Asp Pro Ile Ala Ile Thr Asn Gly Asn Val Ser Leu Lys Val Gly Gly Gly Leu Thr Val Glu Gln Asp Ser Gly Asn Leu Lys Val Asn Pro Lys Ala Pro Leu Gln Val Ala Thr Asp Asn Gln Leu Glu Ile Ser Leu Ala Asp Pro Phe Glu Val Lys Asn Lys Lys Leu Ser Leu Lys Val Gly His Gly Leu Lys Val Ile Asp Glu Asn Ile Ser Thr Leu Gln Gly Leu Leu Gly Asn Leu Val Val Leu Thr Gly Met Gly Ile Gly Thr Glu 

Asn Pro Val Ala Thr Thr Val Ser Phe Asn Lys Glu Thr Gly Cys Thr

- Glu Leu Lys Lys Asp Asp Lys Ile Val Gly Ser Ala Val Asn Val Arg 165 170 175
- Leu Gly Gln Asp Gly Gly Leu Thr Phe Asp Lys Lys Gly Asp Leu Val 180 185 190
- Ala Trp Asn Lys Glu Asn Asp Arg Arg Thr Leu Trp Thr Thr Pro Asp 195 200 205
- Pro Ser Pro Asn Cys Lys Val Ser Glu Glu Lys Asp Ser Lys Leu Thr 210 215 220
- Leu Val Leu Thr Lys Cys Gly Ser Gln Ile Leu Ala Ser Val Ser Leu 225 230 235 240
- Leu Val Val Lys Gly Lys Phe Ala Asn Ile Asn Asn Lys Thr Asn Pro 245 250 255
- Gly Glu Asp Tyr Lys Xaa Phe Ser Val Lys Leu Leu Phe Asp Ala Asn 260 265 270
- Gly Lys Leu Leu Thr Gly Ser Ser Leu Asp Gly Asn Tyr Trp Asn Tyr 275 280 285
- Lys Asn Lys Asp Ser Val Ile Gly Ser Pro Tyr Glu Asn Ala Val Pro 290 295 300
- Phe Met Pro Asn Ser Thr Ala Tyr Pro Lys Ile Ile Asn Gly Thr Ala 305 310 315 320
- Asn Pro Glu Asp Lys Lys Ser Ala Ala Lys Lys Thr Ile Val Thr Asn 325 330 335
- Val Tyr Leu Gly Gly Asp Ala Ala Lys Pro Val Ala Thr Thr Ile Ser 340 345 350
- Phe Asn Lys Glu Thr Glu Ser Asn Cys Val Tyr Ser Ile Thr Phe Asp 355 360 365
- Phe Ala Trp Asn Lys Thr Trp Lys Asn Val Pro Phe Asp Ser Ser Ser 370 375 380
- Leu Thr Phe Ser Tyr Ile Ala Gln Glu 385 390

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<211> 353
<212> PRT
<213> Human Adenovirus 51 Fiber Protein
<400> 42
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           5
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Lys Arg Ala Arg Pro Ser Glu Asp Thr Phe Asn Pro Val Tyr Pro Tyr
       20
                    25
Glu Asp Glu Ser Thr Ser Gln His Pro Phe Ile Asn Pro Gly Phe Ile
     35
                  40
                              45
Ser Pro Asn Gly Phe Thr Gln Ser Pro Asp Gly Val Leu Thr Leu Asn
               55
   50
Cys Leu Thr Pro Leu Thr Thr Gly Gly Pro Leu Gln Leu Lys Val
                                       80
             70
                          75
65
Gly Gly Leu Ile Val Asp Asp Thr Asp Gly Thr Leu Gln Glu Asn
          85
                      90
                                   95
Ile Arg Val Thr Ala Pro Ile Thr Lys Asn Asn His Ser Val Glu Leu
                    105
       100
Ser Ile Gly Asn Gly Leu Glu Thr Gln Asn Asn Lys Leu Cys Ala Lys
                  120
                               125
    115
Leu Gly Asn Gly Leu Lys Phe Asn Asn Gly Asp Ile Cys Ile Lys Asp
  130
                135
                             140
Ser Ile Asn Thr Leu Trp Thr Gly Ile Lys Pro Pro Pro Asn Cys Gln
                                        160
             150
                           155
145
Ile Val Glu Asn Thr Asp Thr Asn Asp Gly Lys Leu Thr Leu Val Leu
                       170
                                    175
         165
Val Lys Asn Gly Gly Leu Val Asn Gly Tyr Val Ser Leu Val Gly Val
                                  190
                    185
       180
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Ser Asp Thr Val Asn Gln Met Phe Thr Gln Lys Ser Ala Thr Ile Gln 

210	215	Ser Ser Gly 220	/ Ash Leu Leu	Inr Asp Glu Ser
Asn Leu Lys I 225	le Pro Leu L 230	ys Asn Lys 235	Ser Ser Thr A 240	la Thr Ser Glu
Ala Ala Thr S 245	er Ser Lys A 25		Pro Ser Thr T 255	hr Ala Tyr Pro
Phe Asn Thr 7 260	Γhr Thr Arg 2 265		u Asn Tyr Ile I 70	His Gly Ile Cys
Tyr Tyr Met T 275	Thr Ser Tyr A 280	asp Arg Ser 285		Leu Asn Ile Ser
Ile Met Leu A 290	sn Ser Arg T 295	Thr Ile Ser S	Ser Asn Val Al	la Tyr Ala Ile
Gln Phe Glu 7 305	Ггр Asn Leu 310	Asn Ala Ly 315	ys Glu Ser Pro 320	Glu Ser Asn Ile
Ala Thr Leu T			Phe Ser Tyr I 335	le Ile Glu Asp
Thr Thr Lys C 340	Cys Ile Ser Le 345		Val Ser Thr C	ys Leu Phe Phe
Asn				
<210> 43 <211> 958 <212> PRT <213> Human	n Adenovirus	s 34 Hexon	Protein	
<400> 43 Leu Ser Arg A	Arg Ala Pro (		o Leu Val Lys 15	Met Ala Thr Pro
Ser Met Leu I 20	Pro Gln Trp 2		et His Ile Ala ( 0	Gly Gln Asp Ala
Ser Glu Tyr L 35	eu Ser Pro C 40	Gly Leu Va 45	l Gln Phe Ala	Arg Ala Thr Asp

N. Communication of the Commun

Thr Tyr Val Asn Leu Gly Asn Lys Phe Arg Asn Pro Thr Val Ala Pro 50 55 60
Thr His Asp Val Thr Thr Asp Arg Ser Gln Arg Leu Met Leu Arg Ph 65 70 75 80
Val Pro Val Asp Arg Glu Asp Asn Thr Tyr Ser Tyr Lys Val Arg Tyr 85 90 95
Thr Leu Ala Val Gly Asp Asn Arg Val Leu Asp Met Ala Ser Thr Ph 100 105 110
Phe Asp Ile Arg Gly Val Leu Asp Arg Gly Pro Ser Phe Lys Pro Tyr 115 120 125
Ser Gly Thr Ala Tyr Asn Ser Leu Ala Pro Lys Gly Ala Pro Asn Ala 130 135 140
Ser Gln Trp Leu Asp Lys Gly Val Thr Ser Thr Gly Leu Val Asp Asp 145 150 155 160
Gly Asn Thr Thr Asp Asp Gly Glu Glu Ala Lys Lys Ala Thr Tyr Th 165 170 175
Phe Gly Asn Ala Pro Val Lys Ala Glu Ala Glu Ile Thr Lys Asp Gly 180 185 190
Leu Pro Val Gly Leu Glu Val Ser Thr Glu Gly Pro Lys Pro Ile Tyr 195 200 205
Ala Asp Lys Leu Tyr Gln Pro Glu Pro Gln Val Gly Asp Glu Thr Tr 210 215 220
Thr Asp Leu Asp Gly Lys Thr Glu Glu Tyr Gly Gly Arg Val Leu Ly 225 230 235 240
Pro Glu Thr Lys Met Lys Pro Cys Tyr Gly Ser Phe Ala Lys Pro Thr 245 250 255
Asn Ile Lys Gly Gly Gln Ala Lys Val Lys Pro Lys Glu Asp Asp Gly 260 265 270

Thr Asn Asn Ile Glu Tyr Asp Ile Asp Met Asn Phe Phe Asp Leu Arg 275 280 285

- Ser Gln Arg Ser Glu Leu Lys Pro Lys Ile Val Met Tyr Ala Glu Asn Val Asp Leu Glu Cys Pro Asp Thr His Val Val Tyr Lys Pro Gly Val Ser Asp Ala Ser Ser Glu Thr Asn Leu Gly Gln Gln Ser Met Pro Asn Arg Pro Asn Tyr Ile Gly Phe Arg Asp Asn Phe Ile Gly Leu Met Tyr Tyr Asn Ser Thr Gly Asn Met Gly Val Leu Ala Gly Gln Ala Ser Gln Leu Asn Ala Val Val Asp Leu Gln Asp Arg Asn Thr Glu Leu Ser Tyr Gln Leu Leu Asp Ser Leu Gly Asp Arg Thr Arg Tyr Phe Ser Met Trp Asn Gln Ala Val Asp Ser Tyr Asp Pro Asp Val Arg Val Ile Glu Asn His Gly Val Glu Asp Glu Leu Pro Asn Tyr Cys Phe Pro Leu Asp Gly
- Val Gly Pro Arg Thr Asp Ser Tyr Lys Glu Ile Lys Pro Asn Gly Asp
- Gln Ser Thr Trp Thr Asn Val Asp Pro Thr Gly Ser Ser Glu Leu Ala
- Lys Gly Asn Pro Phe Ala Met Glu Ile Asn Leu Gln Ala Asn Leu Trp
- Arg Ser Phe Leu Tyr Ser Asn Val Ala Leu Tyr Leu Pro Asp Ser Tyr
- Lys Tyr Thr Pro Ser Asn Val Thr Leu Pro Glu Asn Lys Asn Thr Tyr
- Asp Tyr Met Asn Gly Arg Val Val Pro Pro Ser Leu Val Asp Thr Tyr

Val Asn Ile Gly Ala Arg Trp Ser Leu Asp Ala Met Asp Asn Val Asn 530 535 540
Pro Phe Asn His His Arg Asn Ala Gly Leu Arg Tyr Arg Ser Met Leu 545 550 555 560
Leu Gly Asn Gly Arg Tyr Val Pro Phe His Ile Gln Val Pro Gln Lys 565 570 575
Phe Phe Ala Val Lys Asn Leu Leu Leu Leu Pro Gly Ser Tyr Thr Tyr 580 585 590
Glu Trp Asn Phe Arg Lys Asp Val Asn Met Val Leu Gln Ser Ser Leu 595 600 605
Gly Asn Asp Leu Arg Val Asp Gly Ala Ser Ile Ser Phe Thr Ser Ile 610 615 620
Asn Leu Tyr Ala Thr Phe Phe Pro Met Ala His Asn Thr Ala Ser Thr 625 630 635 640
Leu Glu Ala Met Leu Arg Asn Asp Thr Asn Asp Gln Ser Phe Asn Asp 645 650 655
Tyr Leu Ser Ala Ala Asn Met Leu Tyr Pro Ile Pro Ala Asn Ala Thr 660 665 670
Asn Ile Pro Ile Ser Ile Pro Ser Arg Asn Trp Ala Ala Phe Arg Gly 675 680 685
Trp Ser Phe Thr Arg Leu Lys Thr Lys Glu Thr Pro Ser Leu Gly Ser 690 695 700
Gly Phe Asp Pro Tyr Phe Val Tyr Ser Gly Ser Ile Pro Leu Asp Gly 705 710 715 720
Thr Phe Tyr Leu Asn His Thr Phe Lys Lys Val Ser Ile Met Phe Asp 725 730 735
Ser Ser Val Ser Trp Pro Gly Asn Asp Arg Leu Leu Ser Pro Asn Glu 740 745 750
Phe Glu Ile Lys Arg Thr Val Asp Gly Glu Gly Tyr Asn Val Ala Gln 755 760 765

Cys Asn Met Thr Asp Trp Phe Leu Val Gln Met Leu Ala Asn Tyr Asn Ile Gly Tyr Gln Gly Phe Tyr Ile Pro Glu Gly Tyr Lys Asp Arg Met Tyr Ser Phe Phe Arg Asn Phe Gln Pro Met Ser Arg Gln Val Val Asp Glu Val Asn Lys Tyr Asp Phe Lys Ala Val Ile Pro Tyr Gln His Asn Asn Ser Gly Phe Val Gly Tyr Met Ala Pro Thr Met Arg Gln Gly Gln Tyr Pro Ala Asn Tyr Pro Tyr Pro Leu Ile Gly Thr Thr Ala Val Asn Ser Val Thr Gln Lys Lys Phe Leu Cys Asp Arg Thr Met Trp Arg Ile Pro Phe Ser Ser Asn Phe Met Ser Met Gly Ala Leu Thr Asp Leu Gly Gln Asn Met Leu Tyr Ala Asn Ser Ala His Ala Leu Asp Met Thr Phe Glu Val Asp Pro Met Asp Glu Pro Thr Leu Leu Tyr Leu Leu Phe Glu Val Phe Asp Val Val Arg Val Gln Pro His Arg Gly Ile Ile Glu Ala Val Tyr Leu Arg Thr Pro Phe Ser Ala Gly Asn Ala Thr Thr <210>44 <211>946 <212> PRT <213> Human Adenovirus 35 Hexon Protein <400> 44 Leu Ser Arg Arg Ala Pro Gly Phe Pro Leu Val Lys Met Ala Thr Pro 

Ser Met Leu Pro 20	Gln Trp Ala ' 25	Tyr Met His	Ile Ala Gly Gln Asp Ala
Ser Glu Tyr Leu 35	Ser Pro Gly I 40	æu Val Gln I 45	Phe Ala Arg Ala Thr Asp
	Leu Gly Asn 55	Lys Phe Arg	Asn Pro Thr Val Ala Pro
Thr His Asp Val 65 70			Arg Leu Met Leu Arg Phe
Val Pro Val Asp 85	Arg Glu Asp 90	Asn Thr Tyr 95	Ser Tyr Lys Val Arg Tyr
Thr Leu Ala Val 100	Gly Asp Asn 105	Arg Val Leu 110	Asp Met Ala Ser Thr Phe
Phe Asp Ile Arg 115	Gly Val Leu 2 120	Asp Arg Gly 125	Pro Ser Phe Lys Pro Tyr
Ser Gly Thr Ala 130	Tyr Asn Ser I 135	Leu Ala Pro l 140	Lys Gly Ala Pro Asn Ala
Ser Gln Trp Leu 145 15		Val Thr Ser 55	Thr Gly Leu Val Asp Asp 160
Gly Asn Thr Asp 165	o Asp Gly Glu 170	ı Glu Ala Ly 175	s Lys Ala Thr Tyr Thr Phe
	Val Lys Ala 185		Ile Thr Lys Asp Gly Leu
Pro Val Gly Leu 195	Glu Val Ser	Thr Glu Gly 205	Pro Lys Pro Ile Tyr Ala
Asp Lys Leu Ty	r Gln Pro Glu 215	Pro Gln Val 220	Gly Asp Thr Trp Thr Asp
		ı Tyr Gly Gly 235	Arg Val Leu Lys Pro Glu 240
Thr Lys Met Lys	s Pro Cys Tyr 250	Gly Ser Phe	Ala Lys Pro Thr Asn Ile

- Lys Gly Gly Gln Ala Lys Val Lys Pro Lys Glu Asp Asp Gly Thr Asn Asn Ile Tyr Asp Ile Asp Met Asn Phe Phe Asp Leu Arg Ser Gln Arg Ser Glu Leu Lys Pro Lys Ile Val Met Tyr Ala Glu Asn Val Asp Leu Glu Cys Pro Asp Thr His Val Val Tyr Lys Pro Gly Val Ser Asp Ala Ser Ser Glu Thr Asn Leu Gly Gln Gln Met Pro Asn Arg Pro Asn Tyr Ile Gly Phe Arg Asp Asn Phe Ile Gly Leu Met Tyr Tyr Asn Ser Thr Gly Asn Met Gly Val Leu Ala Gly Gln Ala Ser Gln Leu Asn Ala Val Val Asp Leu Gln Asp Arg Asn Thr Glu Leu Ser Tyr Gln Leu Leu Leu Ser Leu Gly Asp Arg Thr Arg Tyr Phe Ser Met Trp Asn Gln Ala Val Asp Ser Tyr Asp Pro Asp Val Arg Val Ile Glu Asn His Gly Val Glu Asp Glu Leu Pro Asn Tyr Cys Phe Pro Leu Asp Gly Val Gly Pro Arg Thr Asp Ser Tyr Lys Glu Ile Pro Asn Gly Asp Gln Ser Thr Trp Thr Asn Val Asp Pro Thr Gly Ser Ser Glu Leu Ala Lys Gly Asn Pro Phe
- Ala Met Glu Ile Asn Leu Gln Ala Asn Leu Trp Arg Ser Phe Leu Tyr 465 470 475 480

Ser Asn Val Ala Leu Tyr Leu Pro Asp Ser Tyr Lys Tyr Thr Ser Asn 485 490 495

Val Thr Leu Pro Glu Asn Lys Asn Thr Tyr Asp Tyr Met Asn Gly Arg 500 505 510
Val Val Pro Pro Ser Leu Val Asp Thr Tyr Val Asn Ile Gly Ala Arg 515 520 525
Trp Ser Leu Asp Ala Met Asp Asn Val Asn Pro Phe Asn His His Arg 530 535 540
Asn Ala Gly Arg Tyr Arg Ser Met Leu Leu Gly Asn Gly Arg Tyr Val 545 550 555 560
Pro Phe His Ile Gln Val Pro Gln Lys Phe Phe Ala Val Lys Asn Leu 565 570 575
Leu Leu Pro Gly Ser Tyr Thr Tyr Glu Trp Asn Phe Arg Lys Asp 580 585 590
Val Asn Met Val Leu Gln Ser Ser Leu Asp Leu Arg Val Asp Gly Ala 595 600 605
Ser Ile Ser Phe Thr Ser Ile Asn Leu Tyr Ala Thr Phe Phe Pro Met 610 615 620
Ala His Asn Thr Ala Ser Thr Leu Glu Ala Met Leu Arg Asn Asp Thr 625 630 635 640
Asn Asp Gln Ser Phe Asn Asp Tyr Leu Ser Ala Ala Asn Met Leu Tyr 645 650 655
Pro Ile Ala Asn Ala Thr Asn Ile Pro Ile Ser Ile Pro Ser Arg Asn 660 665 670
Trp Ala Ala Phe Arg Gly Trp Phe Thr Arg Leu Lys Thr Lys Glu Thr 675 680 685
Pro Ser Leu Gly Ser Gly Phe Asp Pro Tyr Phe Val Tyr Ser Gly Ser 690 695 700
Ile Pro Tyr Leu Asp Gly Thr Phe Tyr Leu His Thr His Lys Lys Val 705 710 715 720
Ser Ile Met Phe Asp Ser Ser Val Ser Trp Pro Gly Asn Asp Arg Leu 725 730 735

Leu Ser Pro Asn Glu Phe Glu Ile Lys Arg Thr Val Asp Gly Glu Gly Tyr Asn Val Ala Gln Cys Asn Met Thr Lys Asp Trp Phe Leu Val Trp Leu Ala Asn Tyr Asn Ile Gly Tyr Gln Gly Phe Tyr Ile Pro Glu Gly Tyr Lys Asp Arg Met Tyr Ser Phe Phe Arg Asn Phe Gln Pro Met Ser Arg Gln Val Val Asp Glu Val Asn Tyr Lys Asp Phe Lys Ala Val Ala Ile Pro Tyr Gln His Asn Asn Gly Phe Val Gly Tyr Met Ala Pro Thr Met Arg Gln Gly Gln Pro Tyr Pro Ala Asn Tyr Pro Tyr Pro Leu Ile Gly Thr Thr Ala Val Asn Ser Val Thr Gln Lys Lys Phe Leu Cys Asp Arg Thr Met Trp Arg Ile Pro Phe Ser Ser Asn Phe Met Ser Ala Leu Thr Asp Leu Gly Gln Asn Met Leu Tyr Ala Asn Ser Ala His Ala Leu Asp Met Thr Phe Glu Val Asp Pro Met Asp Glu Pro Thr Leu Leu Tyr Leu Leu Phe Glu Val Phe Asp Val Val Arg Val His Gln Pro His Arg Gly Ile Ile Glu Ala Val Leu Arg Thr Pro Phe Ser Ala Gly Asn Ala Thr Thr 

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Ser Met Leu Pro Gln Trp Ala Tyr Met His Ile Ala Gly Gln Asp Ala 20 25 30
Ser Glu Tyr Leu Ser Pro Gly Leu Val Gln Phe Ala Arg Ala Thr Asp 35 40 45
Thr Tyr Phe Asn Leu Gly Asn Lys Phe Arg Asn Pro Thr Val Ala Pro 50 55 60
Thr His Asp Val Thr Thr Asp Arg Ser Gln Arg Leu Met Leu Arg Phe 65 70 75 80
Val Pro Val Asp Arg Glu Asp Asn Thr Tyr Ser Tyr Lys Val Arg Tyr 85 90 95
Thr Leu Ala Val Gly Asp Asn Arg Val Leu Asp Met Ala Ser Thr Phe 100 105 110
Phe Asp Ile Arg Gly Val Leu Asp Arg Gly Pro Ser Phe Lys Pro Tyr 115 120 125
Ser Gly Thr Ala Tyr Asn Ser Leu Ala Pro Lys Gly Ala Pro Asn Ala 130 135 140
Ser Gln Trp Leu Asp Lys Gly Val Thr Ser Thr Gly Leu Val Asp Asp 145 150 155 160
Gly Asn Thr Asp Asp Gly Glu Glu Ala Lys Lys Ala Thr Tyr Thr Phe 165 170 175
Gly Asn Ala Pro Val Lys Ala Glu Ala Glu Ile Thr Lys Asp Gly Leu 180 185 190
Pro Val Gly Leu Glu Val Ser Thr Glu Gly Pro Lys Pro Ile Tyr Ala 195 200 205

Asp Lys Leu Tyr Gln Pro Glu Pro Gln Val Gly Asp Thr Trp Thr Asp 210 215 220

- Leu Asp Gly Lys Thr Glu Glu Tyr Gly Gly Arg Val Leu Lys Pro Glu Thr Lys Met Lys Pro Cys Tyr Gly Ser Phe Ala Lys Pro Thr Asn Ile Lys Gly Gly Gln Ala Lys Val Lys Pro Lys Glu Asp Asp Gly Thr Asn Asn Ile Tyr Asp Ile Asp Met Asn Phe Phe Asp Leu Arg Ser Gln Arg Ser Glu Leu Lys Pro Lys Ile Val Met Tyr Ala Glu Asn Val Asp Leu Glu Cys Pro Asp Thr His Val Val Tyr Lys Pro Gly Val Ser Asp Ala Ser Ser Glu Thr Asn Leu Gly Gln Gln Ser Met Pro Asn Arg Pro Asn Tyr Ile Gly Phe Arg Asp Asn Phe Ile Gly Leu Met Tyr Tyr Asn Ser Thr Gly Asn Met Gly Val Leu Ala Gly Gln Ala Ser Gln Leu Asn Ala Val Val Asp Leu Gln Asp Arg Asn Thr Glu Leu Ser Tyr Gln Leu Leu Asp Ser Leu Gly Asp Arg Thr Arg Tyr Phe Ser Met Trp Asn Gln Ala Val Asp Ser Tyr Asp Pro Asp Val Arg Val Ile Glu Asn His Gly Val Glu Asp Glu Leu Pro Asn Tyr Cys Phe Pro Leu Asp Gly Val Gly Pro
- Thr Asn Val Asp Pro Thr Gly Ser Ser Glu Leu Ala Lys Gly Asn Pro

Arg Thr Asp Ser Tyr Lys Ile Lys Pro Asn Gly Asp Gln Ser Thr Trp 

Phe Ala Met Glu Ile Asn Leu Gln Ala Asn Leu Trp Arg Ser Phe Leu 465 470 475 480
Tyr Ser Asn Val Ala Leu Tyr Leu Pro Asp Ser Tyr Lys Tyr Thr Pro 485 490 495
Ser Asn Val Thr Leu Pro Glu Asn Lys Asn Thr Tyr Asp Tyr Met Asn 500 505 510
Gly Arg Val Val Pro Pro Ser Leu Val Asp Thr Tyr Val Asn Ile Gly 515 520 525
Ala Arg Trp Ser Leu Asp Ala Met Asp Asn Val Asn Pro Phe Asn His 530 535 540
His Arg Ala Gly Leu Arg Tyr Arg Ser Met Leu Leu Gly Asn Gly Arg 545 550 555 560
Tyr Val Pro Phe His Ile Gln Val Pro Gln Lys Phe Phe Ala Val Lys 565 570 575
Asn Leu Leu Leu Pro Gly Ser Tyr Thr Tyr Glu Trp Asn Phe Arg 580 585 590
Lys Asp Val Asn Met Val Leu Gln Ser Leu Gly Asn Asp Leu Arg Val 595 600 605
Asp Gly Ala Ser Ile Ser Phe Thr Ser Ile Asn Leu Tyr Ala Thr Phe 610 615 620
Phe Pro Met Ala His Asn Thr Ala Ser Thr Leu Glu Ala Met Leu Arg 625 630 635 640
Asn Asp Thr Asn Asp Gln Ser Phe Asn Asp Tyr Leu Ser Ala Ala Asn 645 650 655
Met Leu Tyr Pro Ile Pro Ala Asn Ala Thr Asn Ile Pro Ile Ser Ile 660 665 670
Pro Ser Arg Asn Trp Ala Ala Phe Arg Gly Trp Ser Phe Thr Arg Leu 675 680 685
Lys Thr Lys Glu Thr Pro Ser Leu Gly Ser Gly Phe Asp Pro Tyr Phe 690 695 700

- Val Tyr Ser Gly Ser Ile Pro Tyr Asp Gly Thr Phe Tyr Leu Asn His 705 710 715 720
- Thr Phe Lys Lys Val Ser Ile Met Phe Asp Ser Ser Val Ser Trp Pro 725 730 735
- Gly Asn Asp Arg Leu Leu Ser Pro Asn Glu Phe Glu Ile Lys Arg Thr 740 745 750
- Val Asp Gly Asp Gly Tyr Asn Val Ala Gln Cys Asn Met Thr Lys Trp
  755 760 765
- Phe Leu Val Gln Met Leu Ala Asn Tyr Asn Ile Gly Tyr Gln Gly Phe 770 775 780
- Tyr Ile Pro Glu Gly Tyr Lys Asp Arg Met Tyr Ser Phe Phe Arg Asn 785 790 795 800
- Phe Gln Pro Met Ser Arg Gln Val Val Asp Glu Val Asn Tyr Lys Asp 805 810 815
- Phe Lys Ala Val Ile Tyr Gln His Asn Asn Ser Gly Phe Val Gly Tyr 820 825 830
- Met Ala Pro Thr Met Arg Gln Gly Gln Pro Tyr Pro Ala Asn Tyr Pro 835 840 845
- Tyr Pro Leu Ile Gly Thr Thr Ala Val Asn Ser Val Thr Gln Lys Lys 850 855 860
- Phe Leu Cys Asp Arg Thr Met Trp Arg Ile Pro Phe Ser Ser Asn Phe 865 870 875 880
- Met Ser Met Gly Ala Leu Thr Asp Leu Gly Gln Asn Met Leu Tyr Ala 885 890 895
- Asn Ser Ala His Ala Leu Asp Met Thr Phe Glu Val Asp Pro Met Asp 900 905 910
- Glu Pro Thr Leu Leu Tyr Leu Leu Phe Glu Val Phe Asp Val Val Arg 915 920 925
- Val Gln Pro His Arg Gly Ile Ile Glu Ala Val Tyr Leu Arg Thr Pro 930 935 940

Phe Ser Ala Gly Asn Ala Thr Thr 945 950

<210>46

<211>953

<212> PRT

<213> Human Adenovirus 41 Hexon Protein

<400>46

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Arg Phe Pro Leu Val Lys Met Ala Thr Pro Ser Met Met Pro Gln Trp 20 25 30

Ala Tyr Met His Ile Ala Gly Gln Asp Ala Ser Glu Tyr Leu Ser Pro 35 40 45

Gly Leu Val Gln Phe Ala Arg Ala Thr Asp Thr Tyr Phe Ser Leu Gly 50 55 60

Asn Lys Phe Arg Asn Pro Thr Val Ala Pro Thr His Asp Val Thr Thr 65 70 75 80

Asp Arg Ser Gln Arg Leu Thr Leu Arg Phe Ser Pro Ser Asp Arg Glu 85 90 95

Asp Thr Thr Tyr Ser Tyr Lys Ala Arg Phe Thr Leu Ala Gly Asp Asn 100 105 110

Arg Val Leu Asp Met Ala Ser Thr Tyr Phe Asp Ile Arg Gly Val Leu 115 120 125

Asp Arg Gly Pro Ser Phe Lys Pro Tyr Ser Gly Thr Ala Tyr Asn Ser 130 135 140

Leu Ala Pro Lys Gly Ala Pro Asn Ser Ser Gln Trp Ala Asp Lys Glu 145 150 155 160

Arg Val Asn Gly Gly Gly Asn Thr Lys Asp Val Thr Lys Thr Phe Gly 165 170 175

Val Ala Ala Met Gly Gly Glu Asp Ile Thr Glu Lys Gly Leu Lys Ile 180 185 190 Gly Thr Asp Thr Thr Ala Asn Glu Pro Ile Phe Ala Asp Lys Asn Phe Gln Pro Glu Pro Gln Val Gly Glu Glu Asn Gln Glu Thr Phe Val Phe Tyr Gly Gly Arg Ala Leu Lys Lys Glu Thr Lys Met Lys Pro Cys Tyr Gly Ser Phe Ala Arg Pro Thr Asn Glu Lys Gly Gln Ala Lys Phe Ile Ile Gly Asp Asn Gly Gln Pro Thr Glu Asn His Asp Ile Thr Met Ala Phe Asp Thr Pro Gly Gly Thr Ile Thr Gly Gly Thr Gly Gly Pro Gln Asp Glu Leu Lys Ala Asp Ile Val Met Tyr Thr Glu Asn Ile Asn Leu Glu Thr Pro Asp Thr His Val Val Tyr Lys Pro Gly Lys Glu Asp Asp Ser Ser Glu Ile Asn Leu Val Gln Ser Met Pro Asn Arg Pro Asn Tyr Ile Gly Phe Arg Asp Asn Phe Val Gly Leu Met Tyr Tyr Asn Ser Thr Gly Asn Met Gly Val Leu Ala Gly Gln Ala Ser Gln Leu Asn Ala Val Val Asp Leu Gln Asp Arg Asn Thr Glu Leu Ser Tyr Gln Leu Leu Asp Ser Leu Gly Asp Arg Thr Arg Tyr Phe Ser Met Trp Asn Ser Ala Val Asp Ser Tyr Asp Pro Asp Val Arg Ile Ile Glu Asn His Gly Val 

Glu Asp Glu Leu Pro Asn Tyr Cys Phe Pro Leu Asp Gly Ser Gly Thr

Asn Ser Ala Phe Gln Gly Lys Ile Lys Gln Asn Gln Asp Gly Asp Val Asn Asp Asp Trp Glu Lys Asp Asp Lys Val Ser Thr Gln Asn Gln Ile Cys Lys Gly Asn Glu Tyr Ala Met Glu Ile Asn Leu Gln Ala Asn Leu Trp Lys Ser Phe Leu Tyr Ser Asn Val Ala Leu Tyr Leu Asp Ser Tyr Lys Tyr Thr Pro Ala Asn Val Thr Leu Pro Thr Asn Thr Asn Thr Glu Tyr Met Asn Gly Arg Val Val Ala Pro Ser Leu Val Asp Ala Tyr Ile Asn Ile Gly Ala Arg Trp Ser Leu Asp Pro Met Asp Asn Val Asn Pro Phe Asn His Arg Asn Ala Gly Leu Arg Tyr Arg Ser Asn Ala Ser Gly Gln Arg Pro Leu Arg Ala Leu Pro His Pro Ser Ala Pro Lys Val Leu 570 · Cys His Gln Glu Pro Ala Pro Ala Pro Gly Leu Leu His Leu Arg Val Glu Leu Pro Gln Gly Arg Gln His Asp Ala Glu Phe Pro Arg Lys Arg Pro Ala Arg Arg Arg Leu Arg Ala Leu Arg Gln Arg Gln Pro Leu Cys His Ile Leu Pro His Gly Ala Gln His Arg Leu His Pro Gly Ser His Ala Ala Gln Arg His Gln Arg Pro Val Leu Gln Arg Leu Pro Leu 

Arg Gln His Ala Leu Pro His Pro Gly Gln Gly His Gln Arg Ala His

Leu His Pro Ala Gln Leu Gly Arg Leu Ser Arg Leu Glu Phe His Pro Ala Gln Asp Gln Gly Asn Ser Phe Pro Arg Leu Gly Phe Arg Pro Leu Leu Cys Leu Leu Gly Leu His Pro Leu Pro Arg Arg Asp Leu Leu Pro Gln Pro His Leu Gln Glu Gly Leu His His Val Arg Leu Leu Gly Gln Leu Ala Arg Gln Arg Thr Ala Val Thr Pro Asn Glu Phe Glu Ile Lys Arg Ser Val Asp Gly Glu Gly Tyr Asn Val Ala Gln Cys Met Thr Lys Asp Trp Phe Leu Val Gln Met Leu Ser His Tyr Asn Ile Gly Tyr Gln Gly Phe His Val Pro Glu Gly Tyr Lys Asp Arg Met Tyr Ser Phe Phe Arg Asn Phe Gln Pro Met Ser Arg Gln Val Val Asp Glu Ile Asn Tyr Lys Asp Tyr Ala Val Thr Leu Pro Phe Gln His Asn Asn Ser Gly Phe Thr Gly Tyr Leu Ala Pro Thr Met Arg Gln Gly Gln Pro Tyr Pro Ala Asn Phe Pro Leu Ile Gly Ser Thr Ala Val Pro Ser Val Thr Gln Lys Lys Phe Leu Cys Asp Arg Val Met Trp Arg Ile Pro Phe Ser Ser Asn 

Phe Met Ser Met Gly Ala Leu Thr Asp Leu Gly Gln Asn Met Leu Tyr

Ala Asn Ser Ala His Ala Leu Asp Ile Thr Phe Glu Val Asp Pro Met

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Asp Glu Pro Thr Leu Leu Tyr Leu Leu Phe Glu Val Phe Asp Val Val
                              925
                 920
    915
Val His Gln Pro His Arg Gly Val Ile Glu Ala Val Tyr Leu Arg Thr
                            940
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